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Sequence Listing was accepted.

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Reviewer: Durreshwar Anjum

Timestamp: [year=2008; month=3; day=27; hr=14; min=51; sec=34; ms=841;]

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Application No: 10769831

Version No: 3.0

Input Set:

Output Set:

Started: 2008-03-17 11:28:35.991

Finished: 2008-03-17 11:28:38.017

Elapsed: 0 hr(s) 0 min(s) 2 sec(s) 26 ms

Total Warnings: 22

Total Errors: 0

No. of SeqIDs Defined: 23

Actual SeqID Count: 23

Error code	Error Description
W 402	Undefined organism found in <213> in SEQ ID (1)
W 213	Artificial or Unknown found in <213> in SEQ ID (2)
W 213	Artificial or Unknown found in <213> in SEQ ID (3)
W 213	Artificial or Unknown found in <213> in SEQ ID (4)
W 213	Artificial or Unknown found in <213> in SEQ ID (5)
W 213	Artificial or Unknown found in <213> in SEQ ID (6)
W 213	Artificial or Unknown found in <213> in SEQ ID (7)
W 213	Artificial or Unknown found in <213> in SEQ ID (8)
W 213	Artificial or Unknown found in <213> in SEQ ID (9)
W 213	Artificial or Unknown found in <213> in SEQ ID (10)
W 213	Artificial or Unknown found in <213> in SEQ ID (11)
W 213	Artificial or Unknown found in <213> in SEQ ID (12)
W 213	Artificial or Unknown found in <213> in SEQ ID (13)
W 213	Artificial or Unknown found in <213> in SEQ ID (14)
W 213	Artificial or Unknown found in <213> in SEQ ID (15)
W 213	Artificial or Unknown found in <213> in SEQ ID (16)
W 213	Artificial or Unknown found in <213> in SEQ ID (17)
W 213	Artificial or Unknown found in <213> in SEQ ID (18)
W 213	Artificial or Unknown found in <213> in SEQ ID (19)
W 213	Artificial or Unknown found in <213> in SEQ ID (20)

Input Set:

Output Set:

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Total Warnings: 22
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Error code	Error Description
W 213	Artificial or Unknown found in <213> in SEQ ID (21) This error has occurred more than 20 times, will not be displayed
W 402	Undefined organism found in <213> in SEQ ID (22)

SEQUENCE LISTING

<110> Schwabe, Nikolai F
 Tan, Linda C
 Catherine, Napper E
 Fry, Jeremy W
 Pang, Susan

<120> CHIMERIC MHC PROTEIN AND OLIGOMER THEREOF

<130> S-844-US

<140> 10769831

<141> 2004-02-02

<150> PCT/EP03/09056

<151> 2003-08-14

<160> 23

<170> PatentIn version 3.5

<210> 1

<211> 9

<212> PRT

<213> Epstein-Barr virus

<400> 1

Gly Leu Cys Thr Leu Val Ala Met Leu
 1 5

<210> 2

<211> 38

<212> DNA

<213> Artificial

<220>

<223> Oligonucleotide (forward)

<400> 2

gcacaccat atgatccagc gtactccaaa gattcagg

38

<210> 3

<211> 36

<212> DNA

<213> Artificial

<220>

<223> Oligonucleotide (reverse)

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ctacaaggat cccatgtctc gatccactt aactat

36

<210> 4
<211> 20
<212> DNA
<213> Artificial

<220>
<223> Oligonucleotide (forward)

<400> 4
taatacgact cactataggg 20

<210> 5
<211> 19
<212> DNA
<213> Artificial

<220>
<223> Oligonucleotide (reverse)

<400> 5
gctagttatt gctcagcgg 19

<210> 6
<211> 15
<212> PRT
<213> Artificial

<220>
<223> Synthetic Construct

<400> 6

Ser Leu Asn Asp Ile Phe Glu Ala Gln Lys Ile Glu Trp His Glu
1 5 10 15

<210> 7
<211> 15
<212> PRT
<213> Artificial

<220>
<223> Synthetic Construct

<400> 7

Pro Gln Pro Gln Pro Lys Pro Gln Pro Lys Pro Glu Pro Glu Thr
1 5 10 15

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<211> 80
<212> DNA
<213> Artificial

<220>
 <223> Oligonucleotide (forward)

<400> 8
 taaagcttca gggccagagc ccgttgggct cagacctggg cccgcagatg cttcgggaac 60
 tgcaggaaac caacgcggcg 80

<210> 9
 <211> 81
 <212> DNA
 <213> Artificial

<220>
 <223> Oligonucleotide (reverse)

<400> 9
 gaacgtgatc tccctgacct gctgccgcag cagctcccgc acgtcctgca gcgccgcgtt 60
 ggtttcctgc agttcccgaa g 81

<210> 10
 <211> 81
 <212> DNA
 <213> Artificial

<220>
 <223> Oligonucleotide (forward)

<400> 10
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 acggtgatgg agtgtgacgc g 81

<210> 11
 <211> 80
 <212> DNA
 <213> Artificial

<220>
 <223> Oligonucleotide (reverse)

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 tccatcaccg tgtttttcag 80

<210> 12
 <211> 108
 <212> DNA
 <213> Artificial

<220>
 <223> Oligonucleotide (forward)

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 ccgaaaccgc agccgaaacc ggaaccggaa actagtttga acgacatc 108

<210> 13
 <211> 96
 <212> DNA
 <213> Artificial

<220>
 <223> Oligonucleotide (reverse)

<400> 13
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 gggtccgggtt tcggctgcgg ttccggctgc ggctgc 96

<210> 14
 <211> 72
 <212> DNA
 <213> Artificial

<220>
 <223> Oligonucleotide (forward)

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 gttctggtgg ta 72

<210> 15
 <211> 72
 <212> DNA
 <213> Artificial

<220>
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<400> 15
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 ccaccaccac cg 72

<210> 16
 <211> 25
 <212> PRT
 <213> Artificial

<220>

<223> Synthetic Construct

<400> 16

Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly
1 5 10 15

Ser Gly Gly Gly Gly Ser Gly Gly Lys
20 25

<210> 17

<211> 24

<212> PRT

<213> Artificial

<220>

<223> Synthetic Construct

<400> 17

Gly Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser
1 5 10 15

Gly Gly Gly Gly Ser Gly Gly Lys
20

<210> 18

<211> 22

<212> DNA

<213> Artificial

<220>

<223> Oligonucleotide (forward)

<400> 18

gagacatggg aggtggtggt gg

22

<210> 19

<211> 22

<212> DNA

<213> Artificial

<220>

<223> Oligonucleotide (reverse)

<400> 19

ccaccaccac ctcccatgtc tc

22

<210> 20

<211> 35

<212> DNA

<213> Artificial

<220>

<223> Oligonucleotide (forward)

<400> 20

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35

<210> 21

<211> 37

<212> DNA

<213> Artificial

<220>

<223> Oligonucleotide (reverse)

<400> 21

gcatacggat ccttacggct cccatctcag ggtgagg

37

<210> 22

<211> 64

<212> PRT

<213> Rat

<400> 22

Gln Gly Gln Ile Pro Leu Gly Gly Asp Leu Ala Pro Gln Met Leu Arg
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Glu Leu Gln Glu Thr Asn Ala Ala Leu Gln Asp Val Arg Glu Leu Leu
20 25 30

Arg Gln Gln Val Lys Glu Ile Thr Phe Leu Lys Asn Thr Val Met Glu
35 40 45

Cys Asp Ala Cys Gly Met Gln Pro Ala Arg Thr Pro Gly Leu Ser Val
50 55 60

<210> 23

<211> 757

<212> PRT

<213> Homo sapiens

<300>

<308> Genbank/1705995

<309> 1996-10-01

<313> (1)..(757)

<400> 23

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Gln Met Leu Arg Glu Leu Gln Glu Thr Asn Ala Ala Leu Gln Asp Val	35	40	45
Arg Asp Trp Leu Arg Gln Gln Val Arg Glu Ile Thr Phe Leu Lys Asn	50	55	60
Thr Val Met Glu Cys Asp Ala Cys Gly Met Gln Gln Ser Val Arg Thr	65	70	75
Gly Leu Pro Ser Val Arg Pro Leu Leu His Cys Ala Pro Gly Phe Cys	85	90	95
Phe Pro Gly Val Ala Cys Ile Gln Thr Glu Ser Gly Gly Arg Cys Gly	100	105	110
Pro Cys Pro Ala Gly Phe Thr Gly Asn Gly Ser His Cys Thr Asp Val	115	120	125
Asn Glu Cys Asn Ala His Pro Cys Phe Pro Arg Val Arg Cys Ile Asn	130	135	140
Thr Ser Pro Gly Phe Arg Cys Glu Ala Cys Pro Pro Gly Tyr Ser Gly	145	150	155
Pro Thr His Gln Gly Val Gly Leu Ala Phe Ala Lys Ala Asn Lys Gln	165	170	175
Val Cys Thr Asp Ile Asn Glu Cys Glu Thr Gly Gln His Asn Cys Val	180	185	190
Pro Asn Ser Val Cys Ile Asn Thr Arg Gly Ser Phe Gln Cys Gly Pro	195	200	205
Cys Gln Pro Gly Phe Val Gly Asp Gln Ala Ser Gly Cys Gln Arg Gly	210	215	220
Ala Gln Arg Phe Cys Pro Asp Gly Ser Pro Ser Glu Cys His Glu His	225	230	235
			240

Ala Asp Cys Val Leu Glu Arg Asp Gly Ser Arg Ser Cys Val Cys Arg
245 250 255

Val Gly Trp Ala Gly Asn Gly Ile Leu Cys Gly Arg Asp Thr Asp Leu
260 265 270

Asp Gly Phe Pro Asp Glu Lys Leu Arg Cys Pro Glu Pro Gln Cys Arg
275 280 285

Lys Asp Asn Cys Val Thr Val Pro Asn Ser Gly Gln Glu Asp Val Asp
290 295 300

Arg Asp Gly Ile Gly Asp Ala Cys Asp Pro Asp Ala Asp Gly Asp Gly
305 310 315 320

Val Pro Asn Glu Lys Asp Asn Cys Pro Leu Val Arg Asn Pro Asp Gln
325 330 335

Arg Asn Thr Asp Glu Asp Lys Trp Gly Asp Ala Cys Asp Asn Cys Arg
340 345 350

Ser Gln Lys Asn Asp Asp Gln Lys Asp Thr Asp Gln Asp Gly Arg Gly
355 360 365

Asp Ala Cys Asp Asp Asp Ile Asp Gly Asp Arg Ile Arg Asn Gln Ala
370 375 380

Asp Asn Cys Pro Arg Val Pro Asn Ser Asp Gln Lys Asp Ser Asp Gly
385 390 395 400

Asp Gly Ile Gly Asp Ala Cys Asp Asn Cys Pro Gln Lys Ser Asn Pro
405 410 415

Asp Gln Ala Asp Val Asp His Asp Phe Val Gly Asp Ala Cys Asp Ser
420 425 430

Asp Gln Asp Gln Asp Gly Asp Gly His Gln Asp Ser Arg Asp Asn Cys
435 440 445

Pro Thr Val Pro Asn Ser Ala Gln Glu Asp Ser Asp His Asp Gly Gln
450 455 460

Gly Asp Ala Cys Asp Asp Asp Asp Asp Asn Asp Gly Val Pro Asp Ser
 465 470 475 480

Arg Asp Asn Cys Arg Leu Val Pro Asn Pro Gly Gln Glu Asp Ala Asp
 485 490 495

Arg Asp Gly Val Gly Asp Val Cys Gln Asp Asp Phe Asp Ala Asp Lys
 500 505 510

Val Val Asp Lys Ile Asp Val Cys Pro Glu Asn Ala Glu Val Thr Leu
 515 520 525

Thr Asp Phe Arg Ala Phe Gln Thr Val Val Leu Asp Pro Glu Gly Asp
 530 535 540

Ala Gln Ile Asp Pro Asn Trp Val Val Leu Asn Gln Gly Arg Glu Ile
 545 550 555 560

Val Gln Thr Met Asn Ser Asp Pro Gly Leu Ala Val Gly Tyr Thr Ala
 565 570 575

Phe Asn Gly Val Asp Phe Glu Gly Thr Phe His Val Asn Thr Val Thr
 580 585 590

Asp Asp Asp Tyr Ala Gly Phe Ile Phe Gly Tyr Gln Asp Ser Ser Ser
 595 600 605

Phe Tyr Val Val Met Trp Lys Gln Met Glu Gln Thr Tyr Trp Gln Ala
 610 615 620

Asn Pro Phe Arg Ala Val Ala Glu Pro Gly Ile Gln Leu Lys Ala Val
 625 630 635 640

Lys Ser Ser Thr Gly Pro Gly Glu Gln Leu Arg Asn Ala Leu Trp His
 645 650 655

Thr Gly Asp Thr Glu Ser Gln Val Arg Leu Leu Trp Lys Asp Pro Arg
 660 665 670

Asn Val Gly Trp Lys Asp Lys Lys Ser Tyr Arg Trp Phe Leu Gln His
 675 680 685

Arg Pro Gln Val Gly Tyr Ile Arg Val Arg Phe Tyr Glu Gly Pro Glu
690 695 700

Leu Val Ala Asp Ser Asn Val Val Leu Asp Thr Thr Met Arg Gly Gly
705 710 715 720

Arg Leu Gly Val Phe Cys Phe Ser Gln Glu Asn Ile Ile Trp Ala Asn
725 730 735

Leu Arg Tyr Arg Cys Asn Asp Thr Ile Pro Glu Asp Tyr Glu Thr His
740 745 750

Gln Leu Arg Gln Ala
755